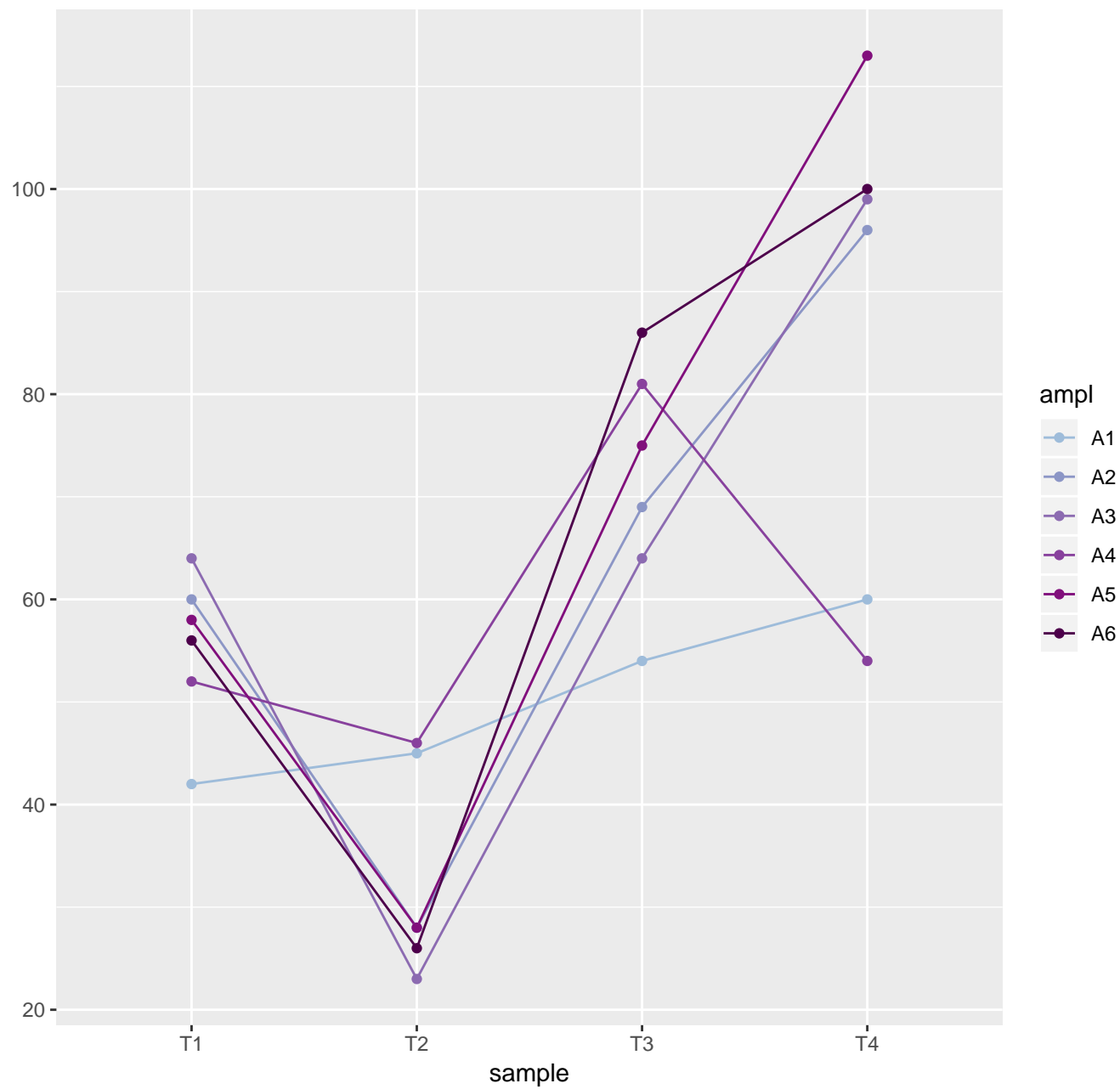
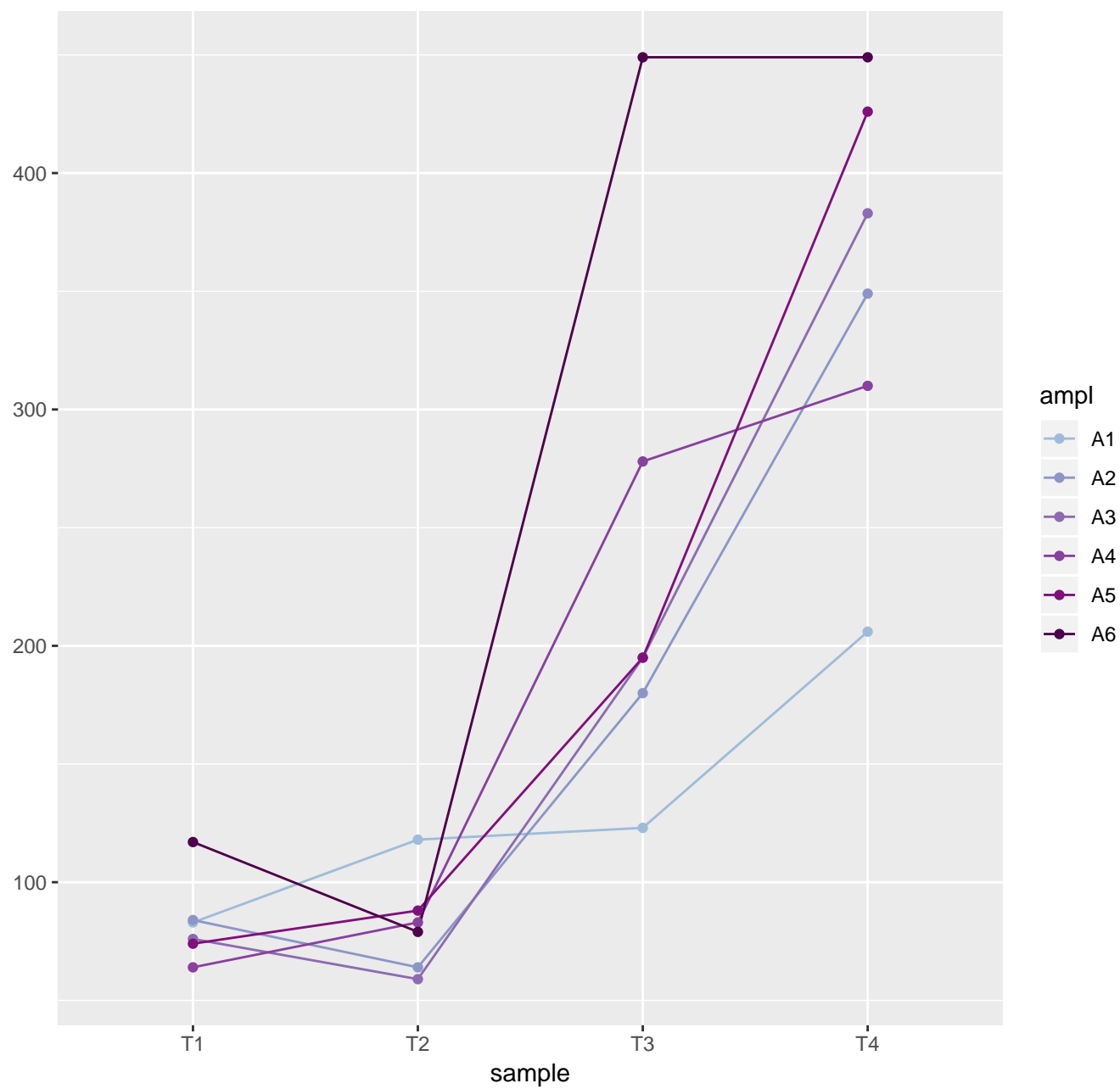


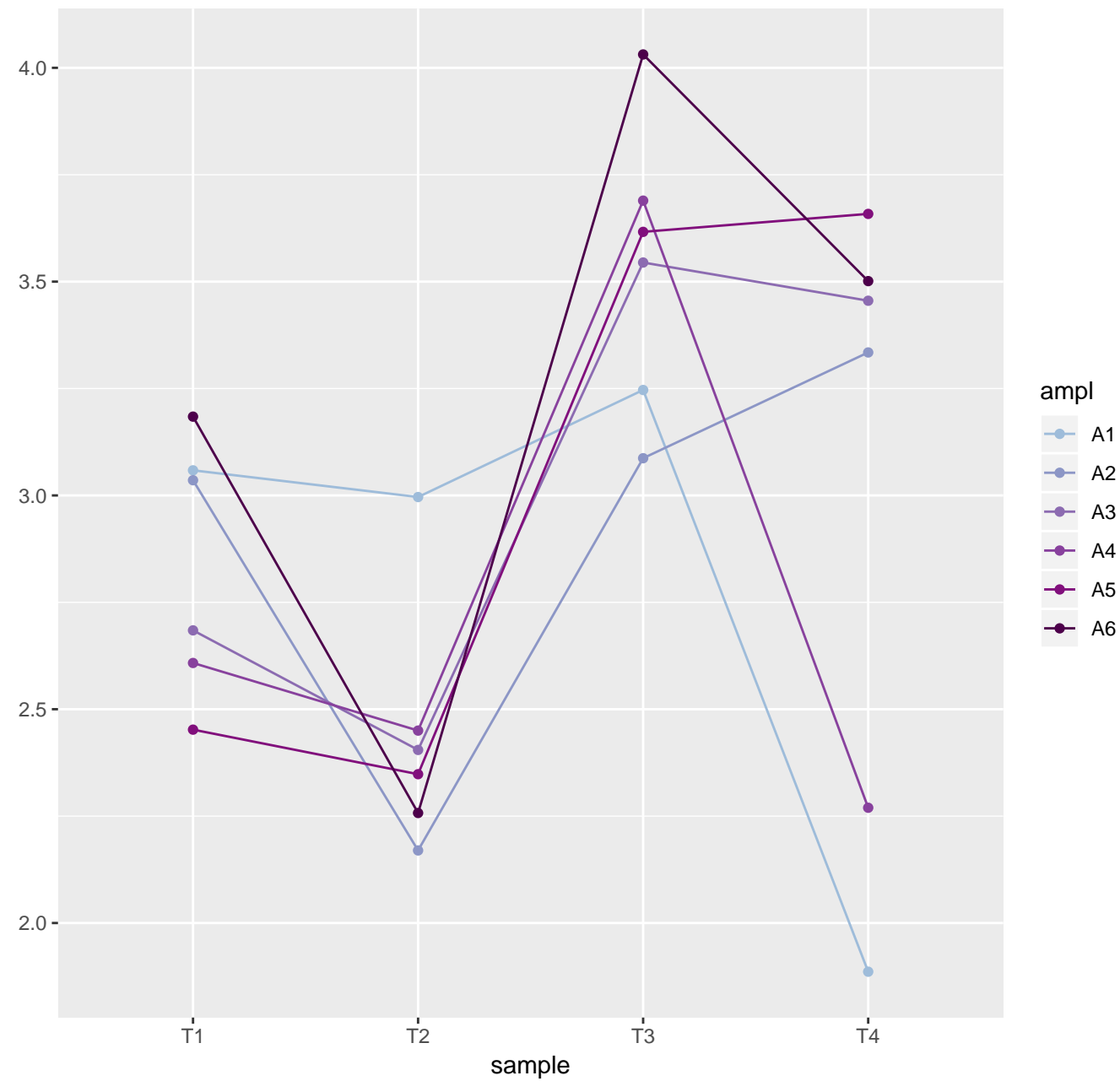
The number of polymorphic sites



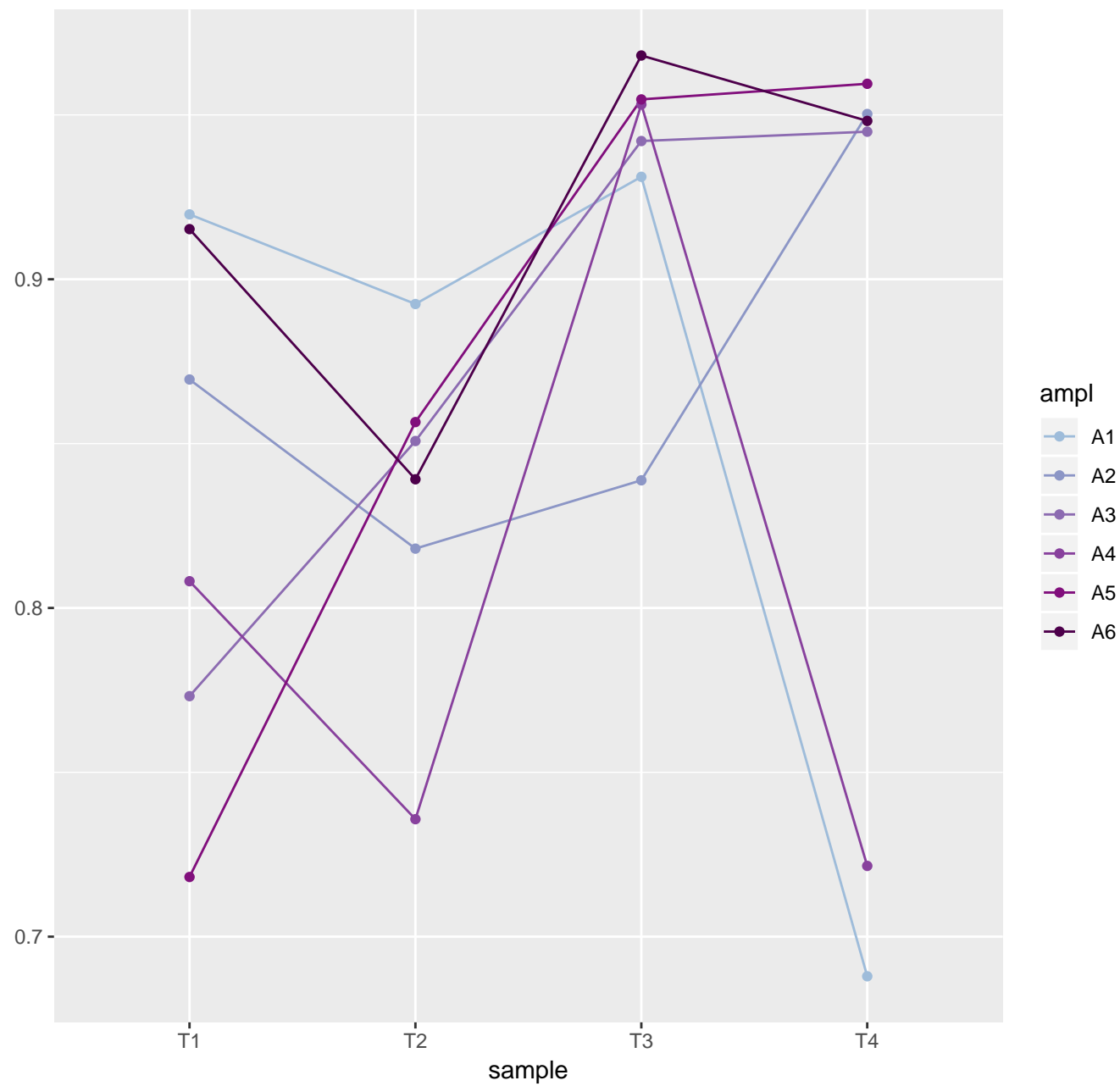
Number of mutations

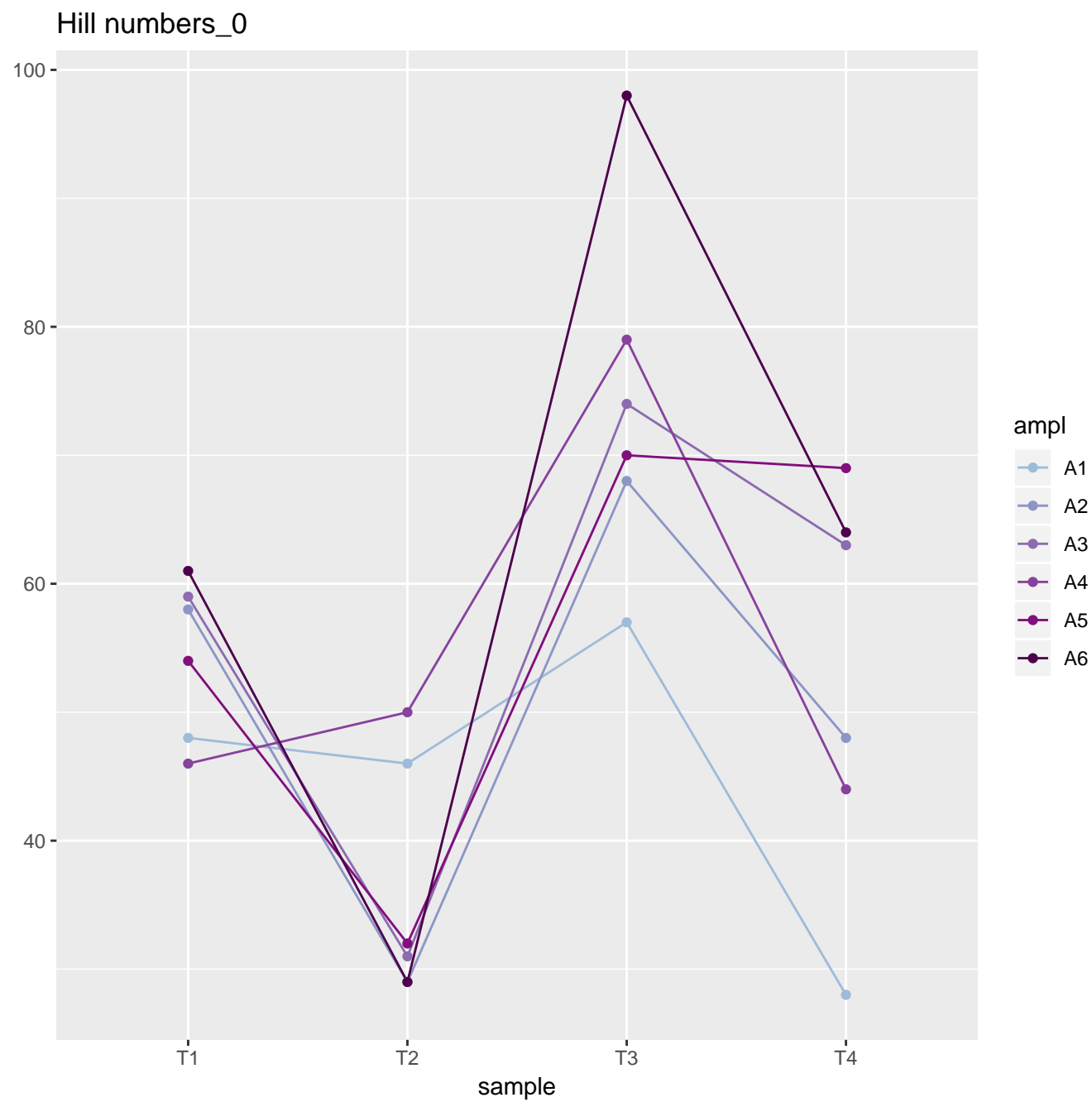


Shannon entropy

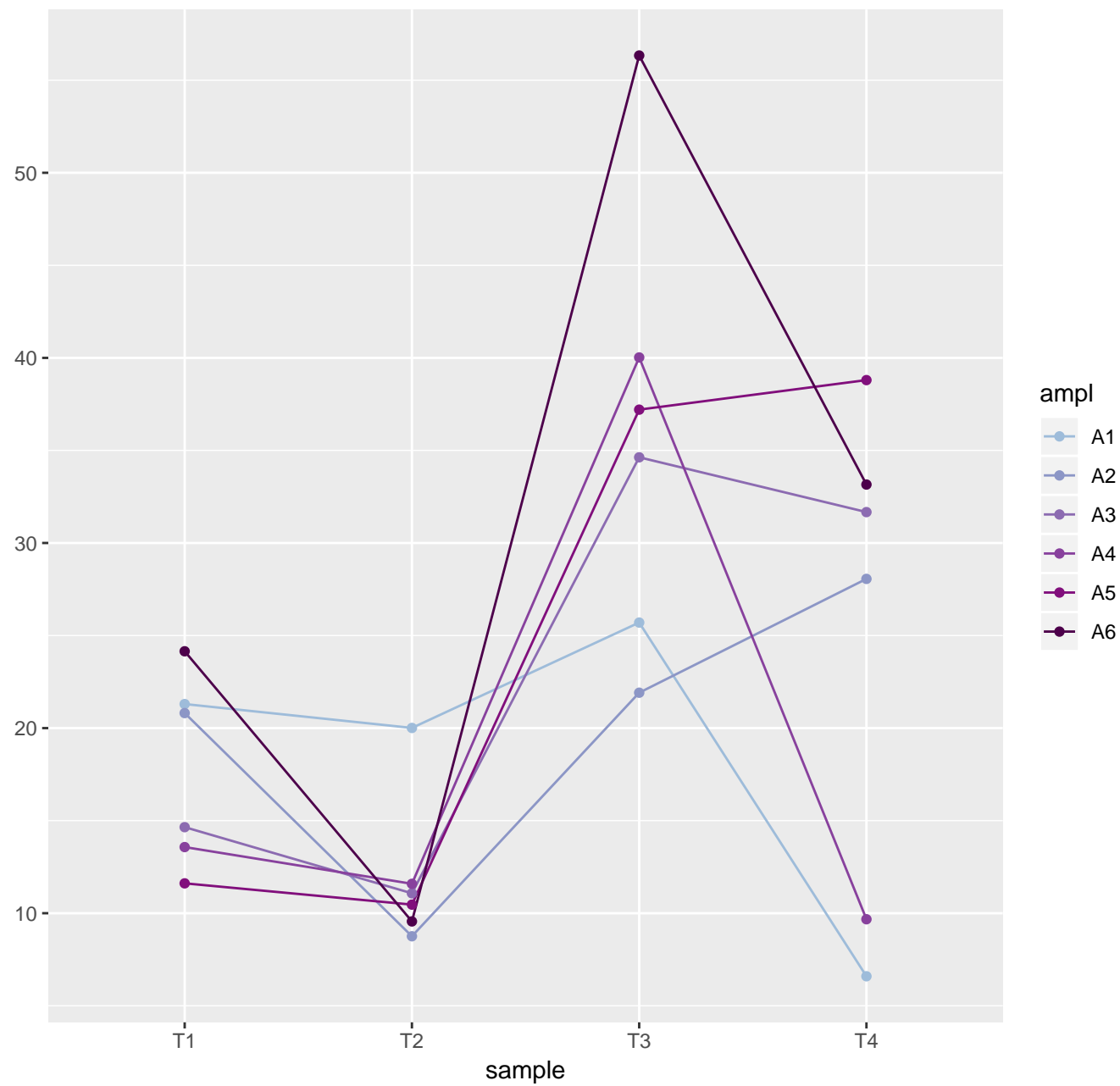


Gini-Simpson

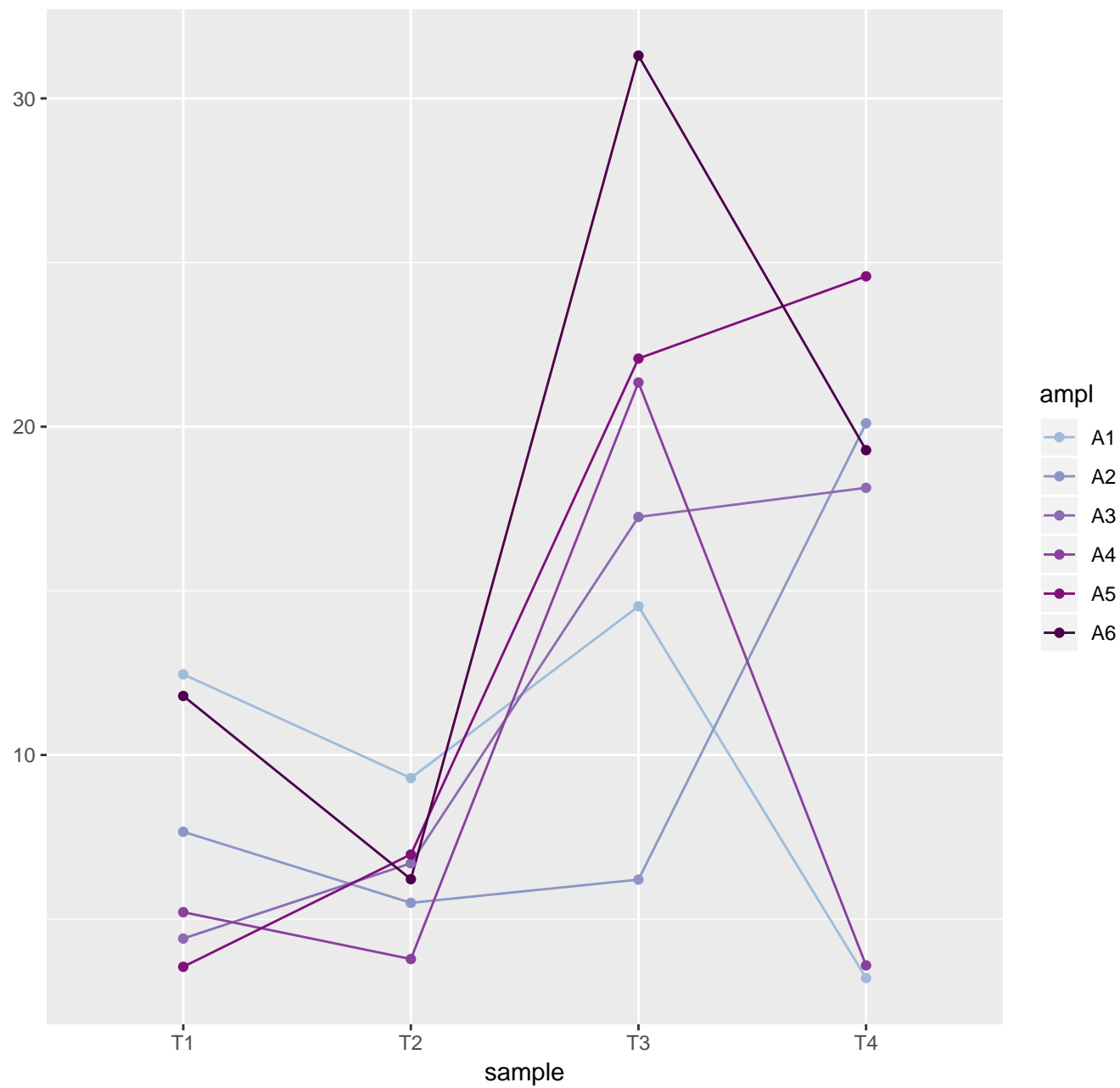




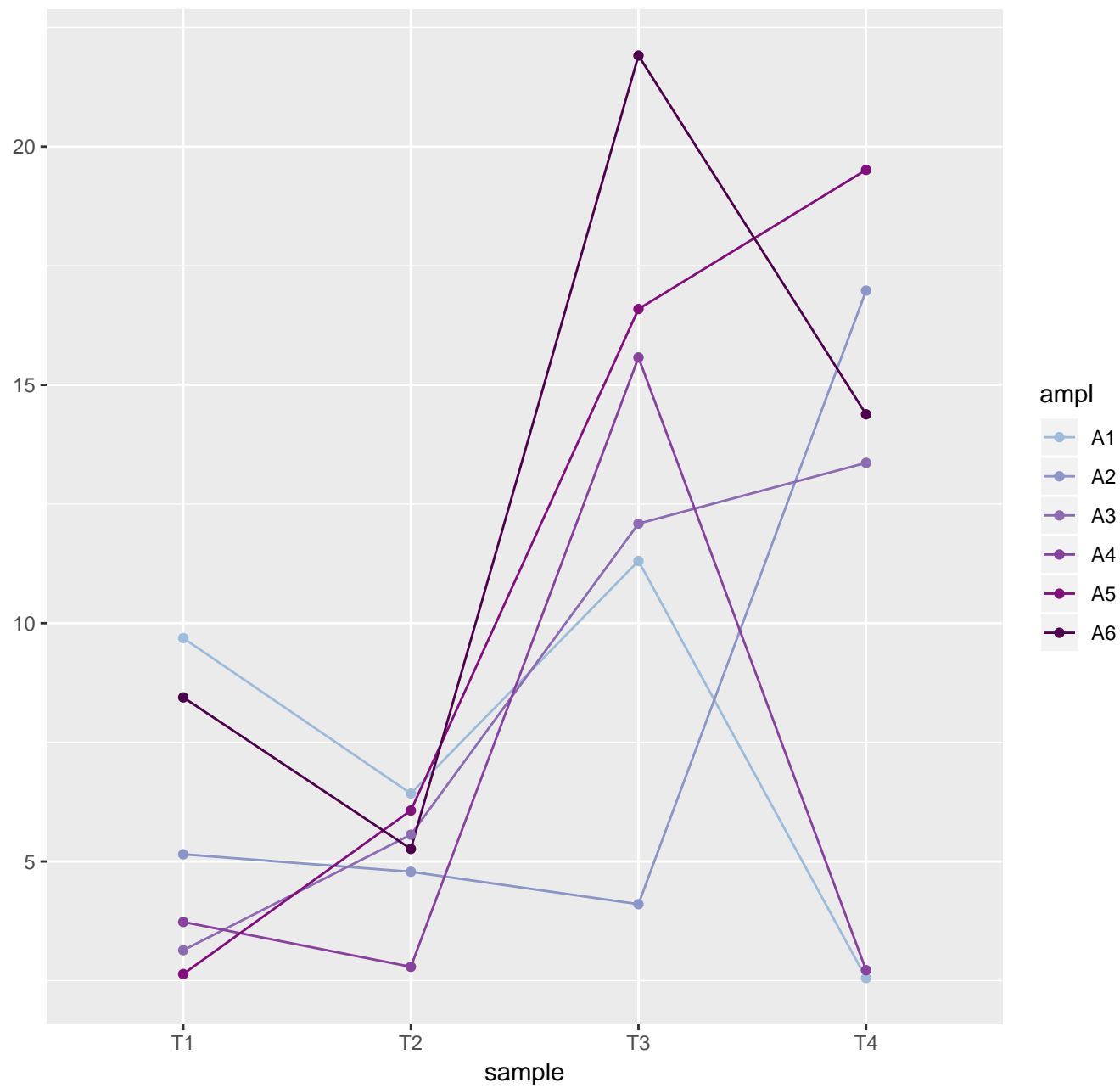
Hill numbers_1



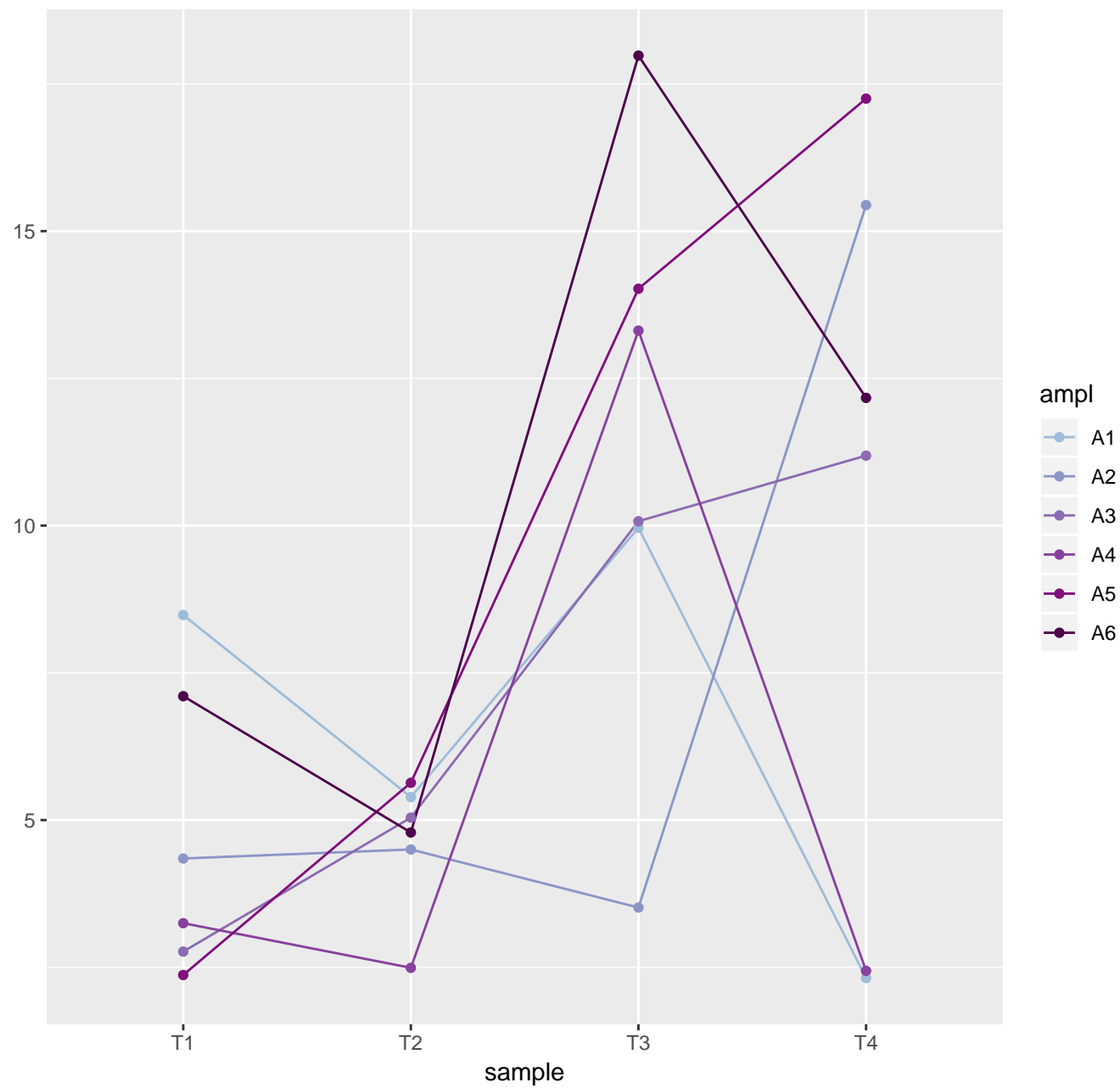
Hill numbers_2



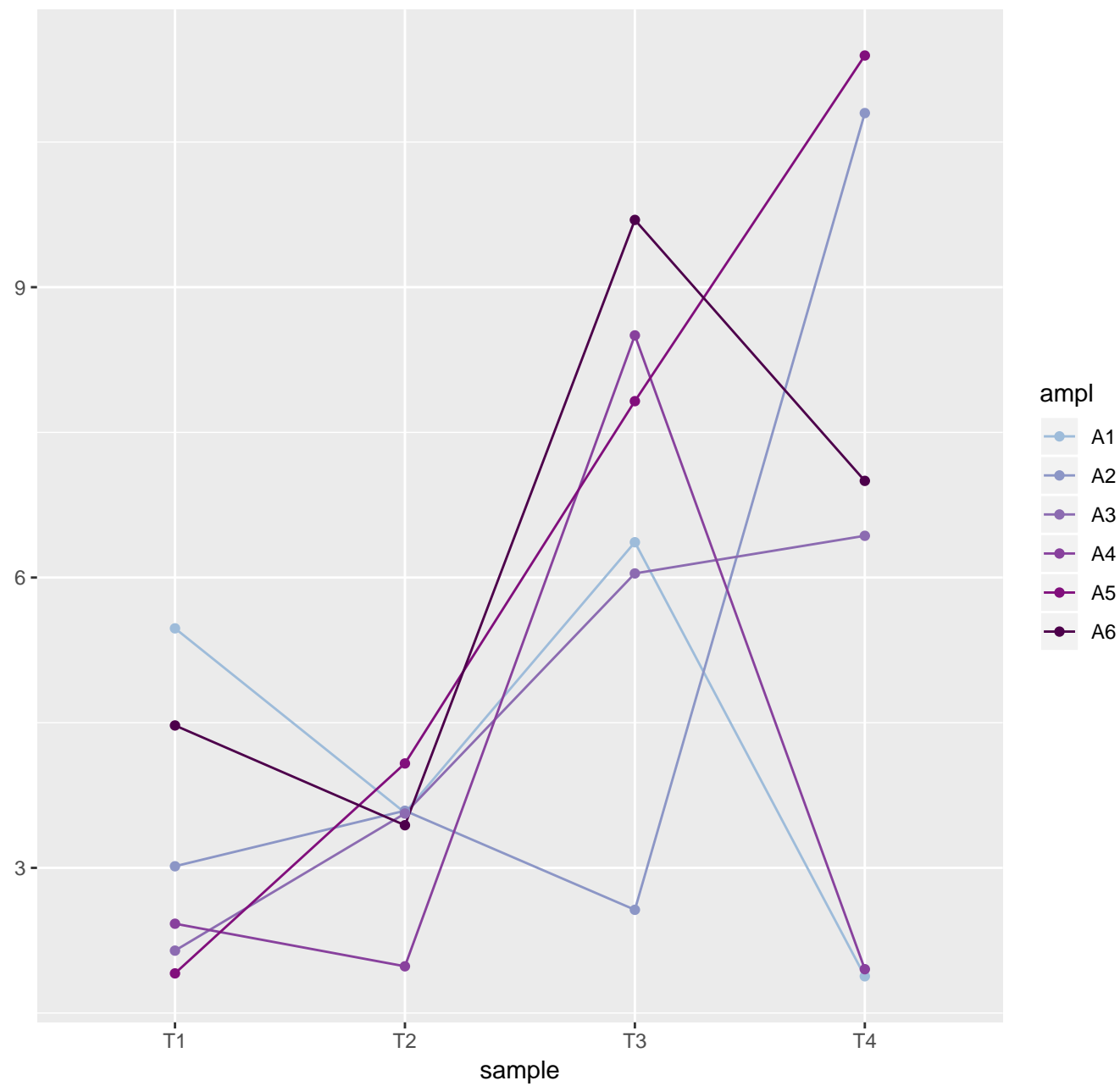
Hill numbers_3



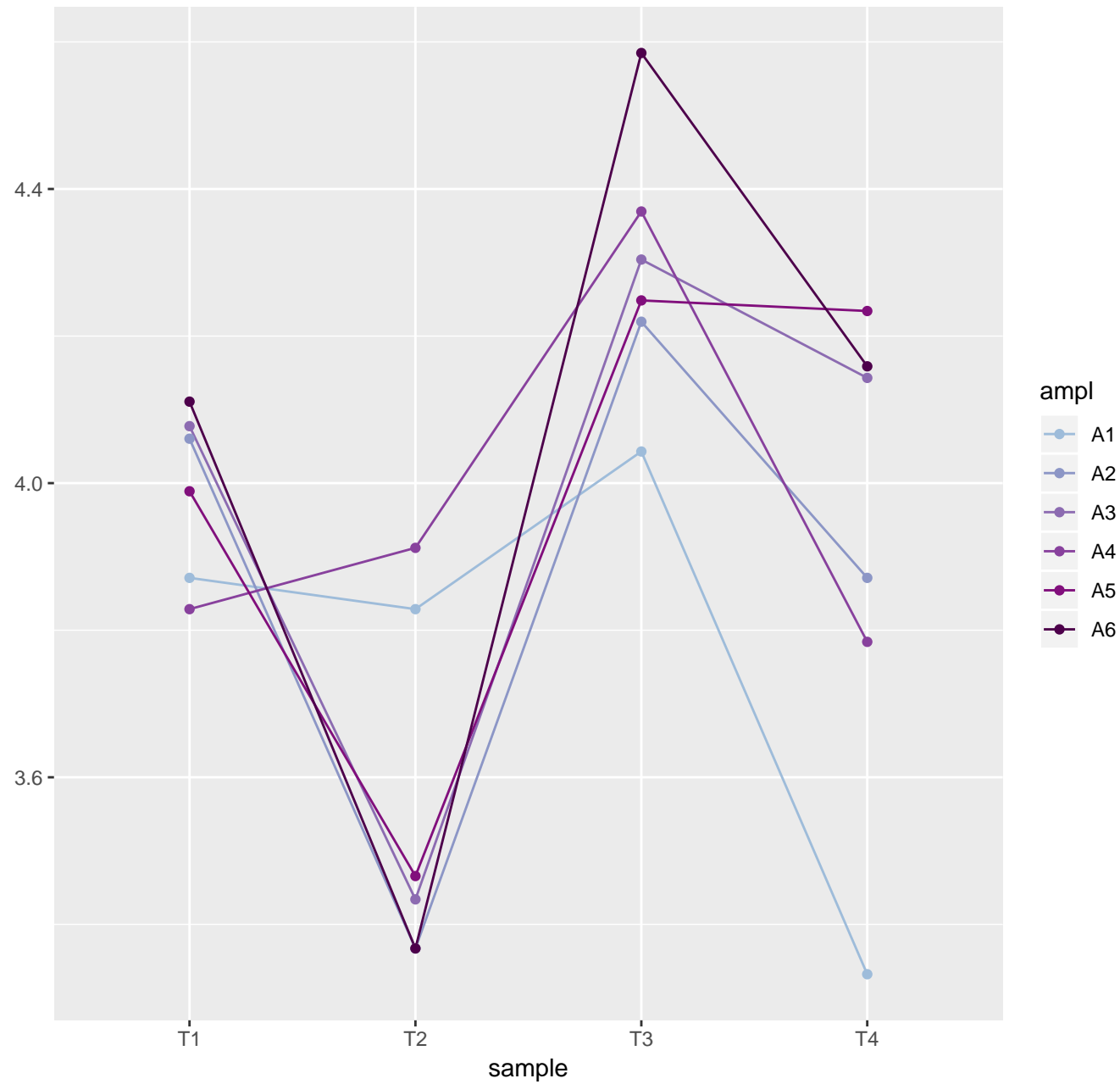
Hill numbers_4



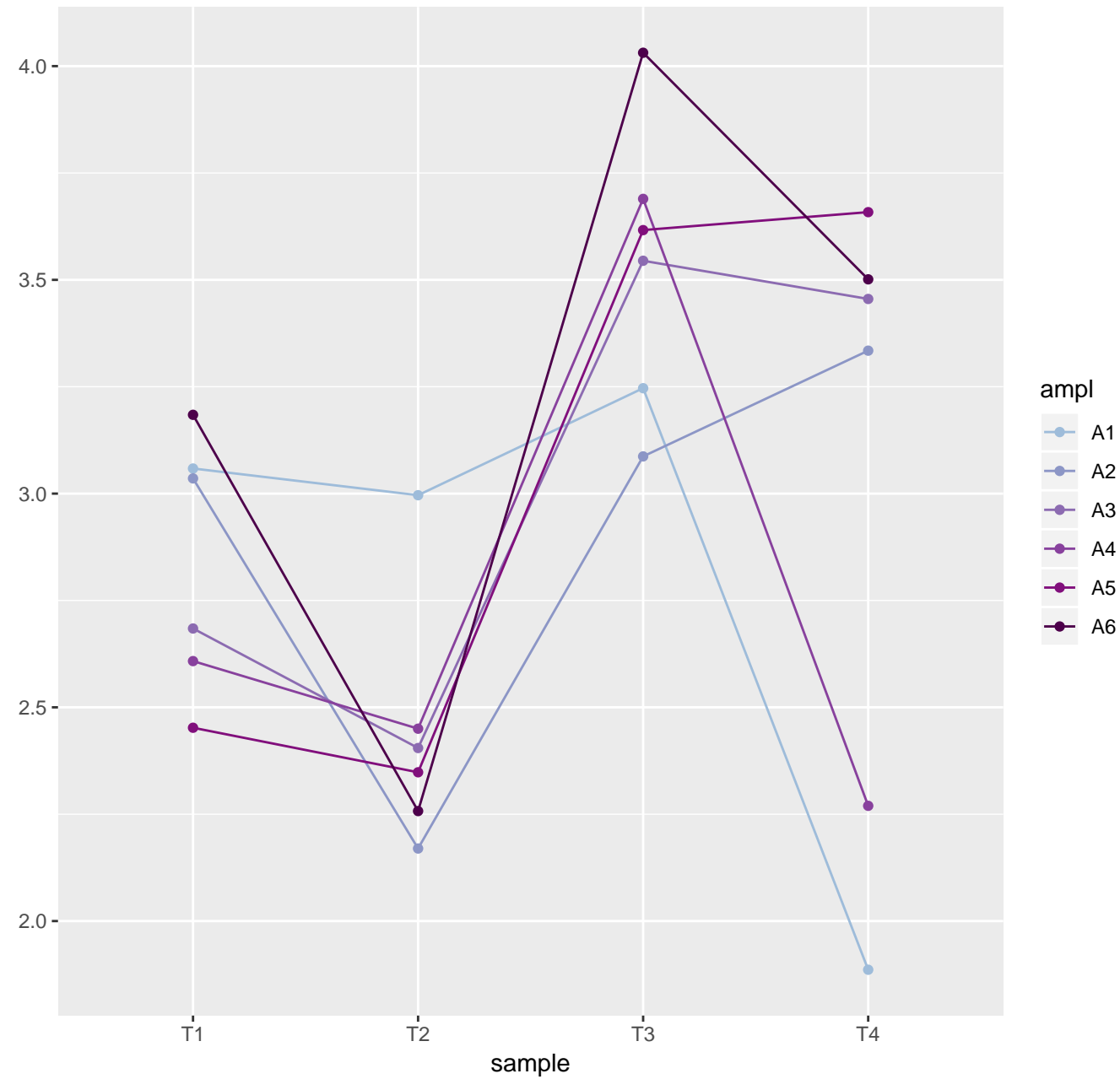
Hill numbers_Inf



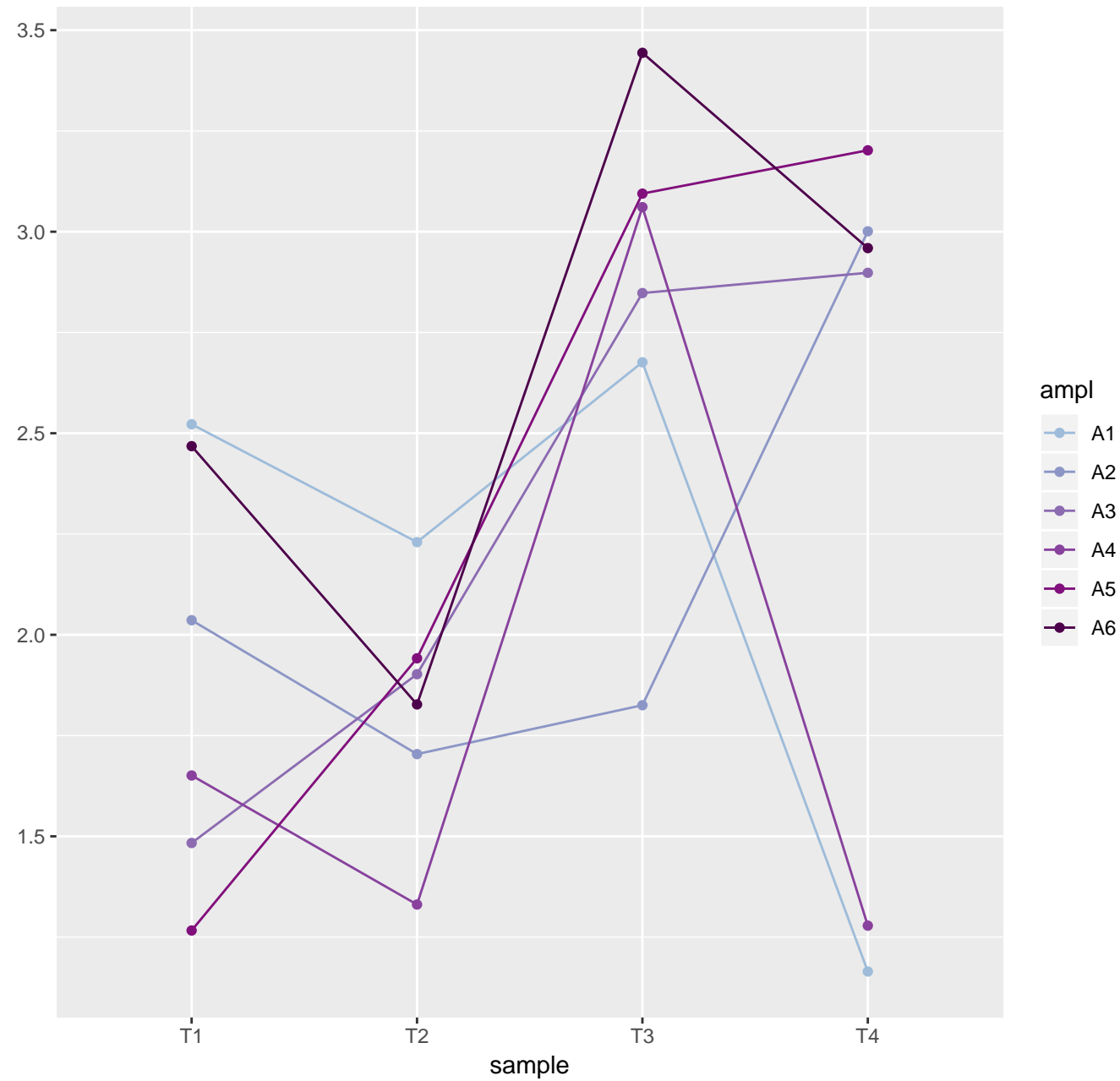
Rényi entropy_0



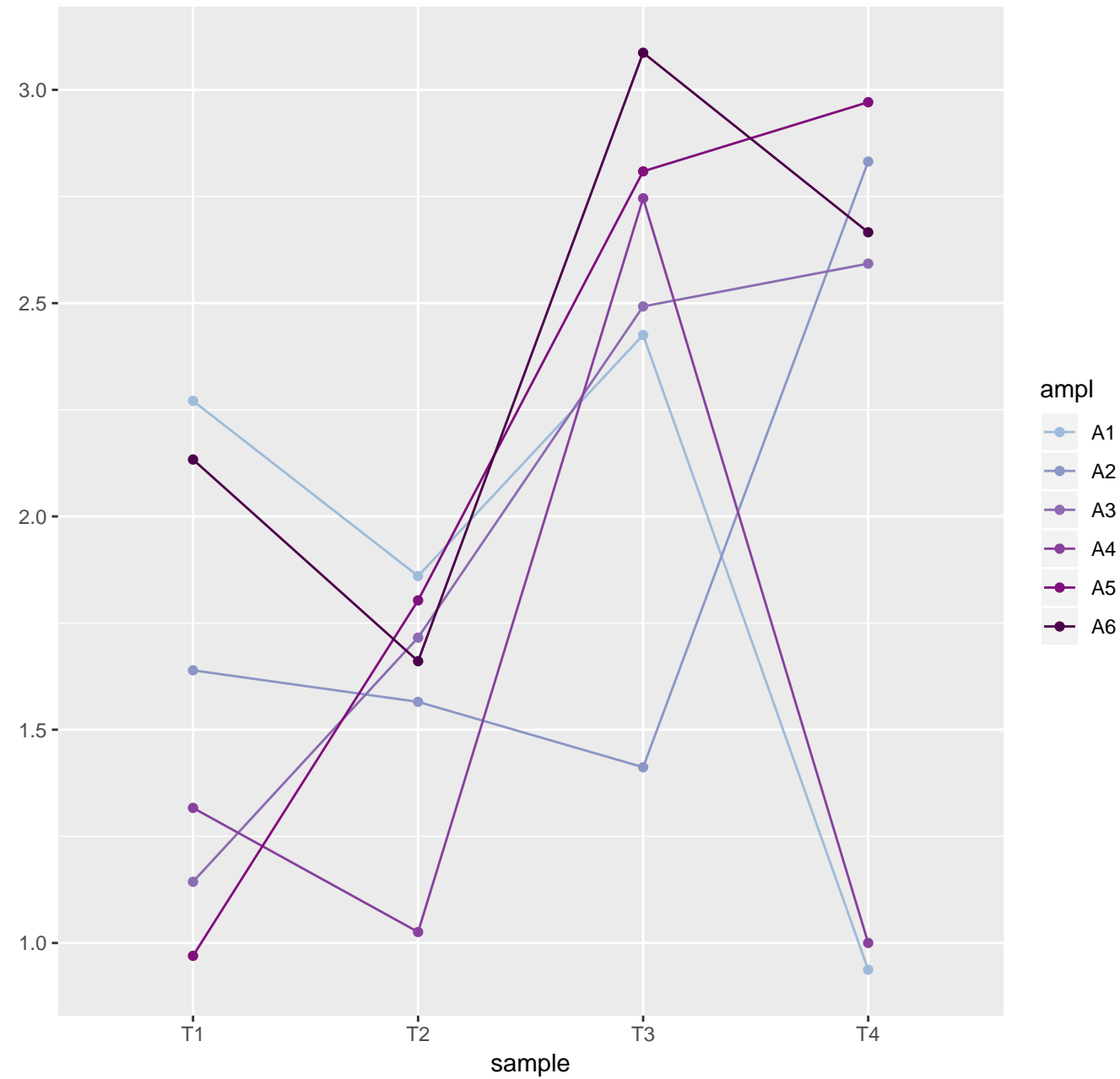
Rényi entropy_1



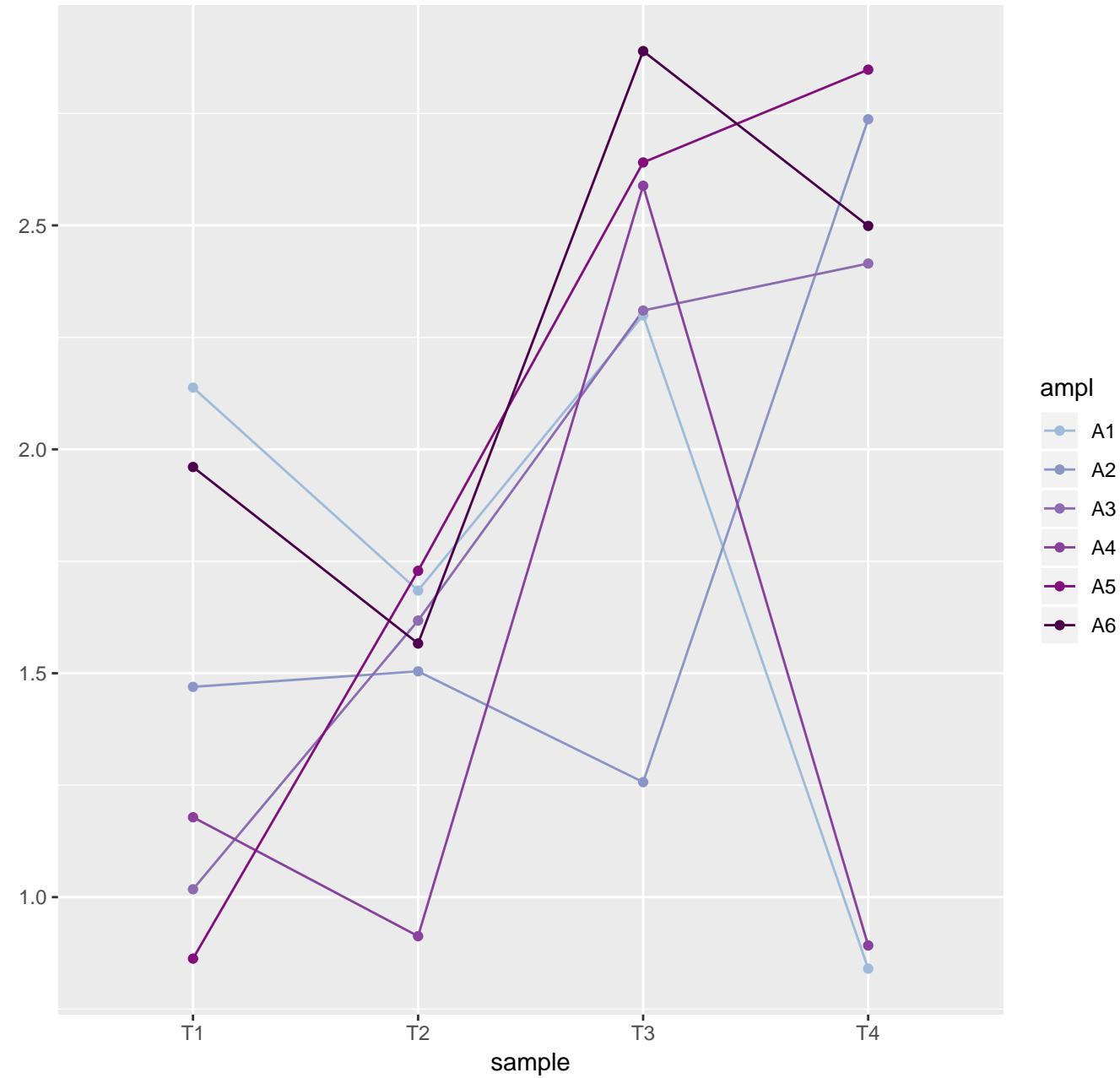
Rényi entropy_2

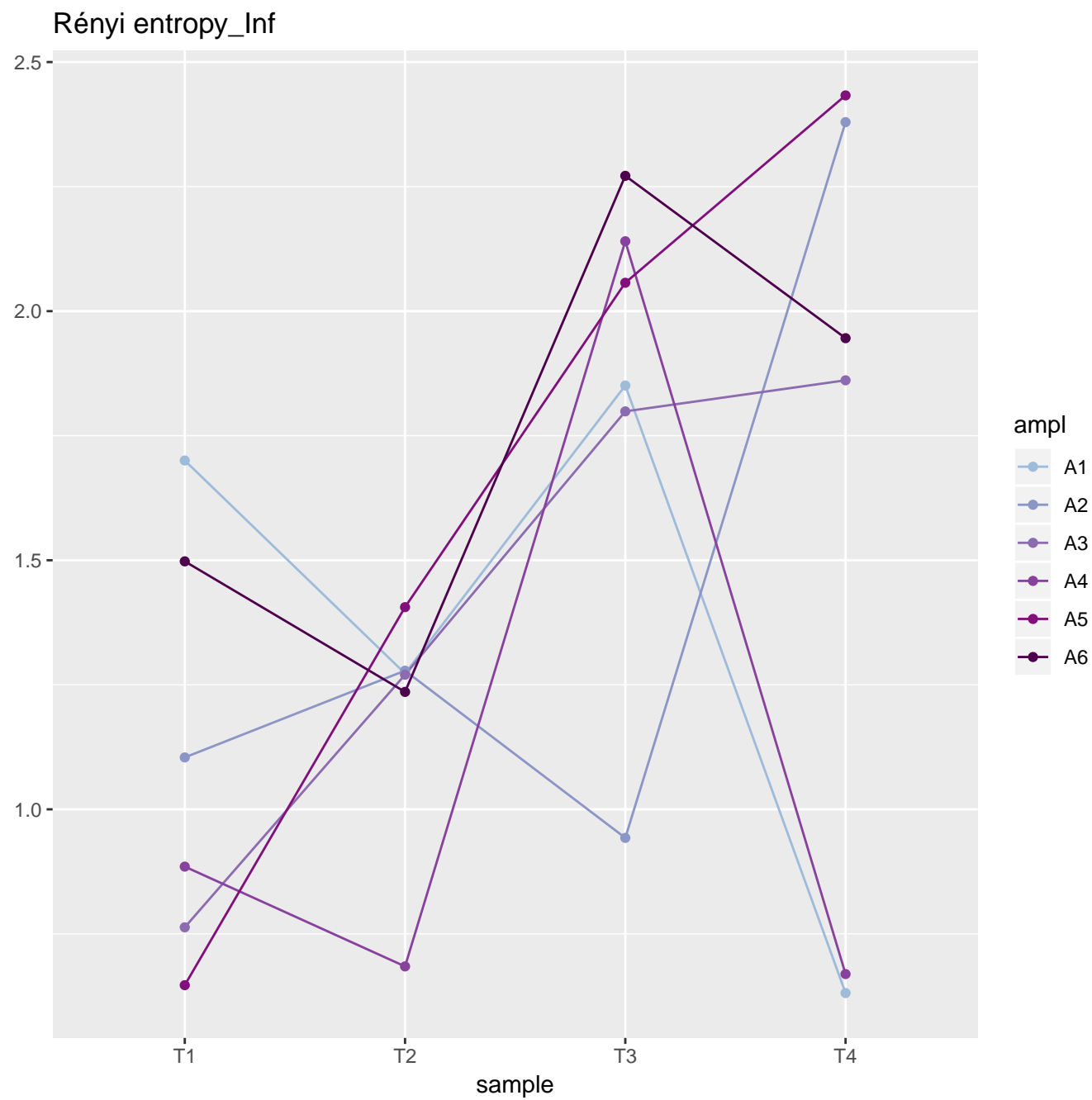


Rényi entropy_3

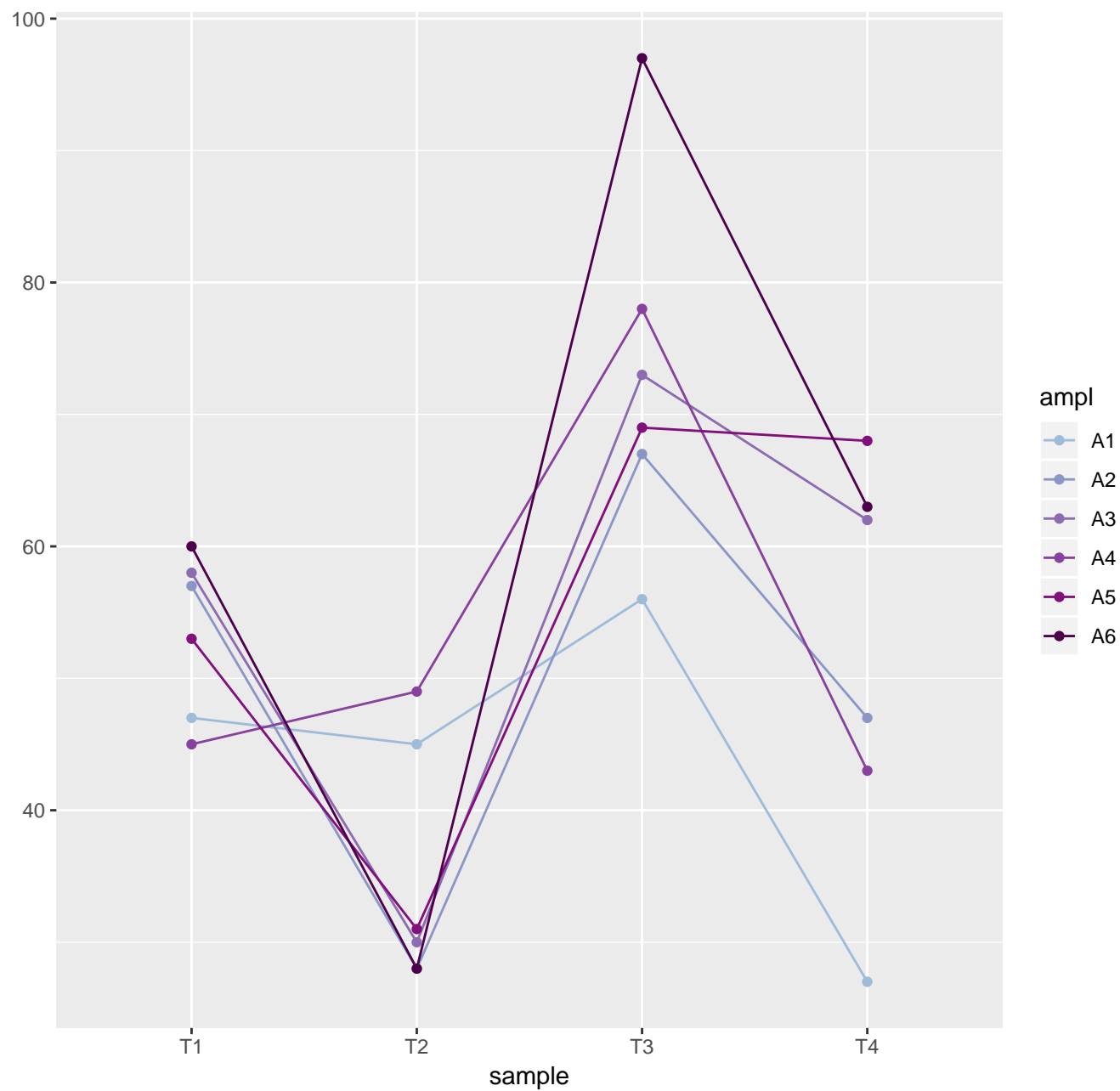


Rényi entropy_4

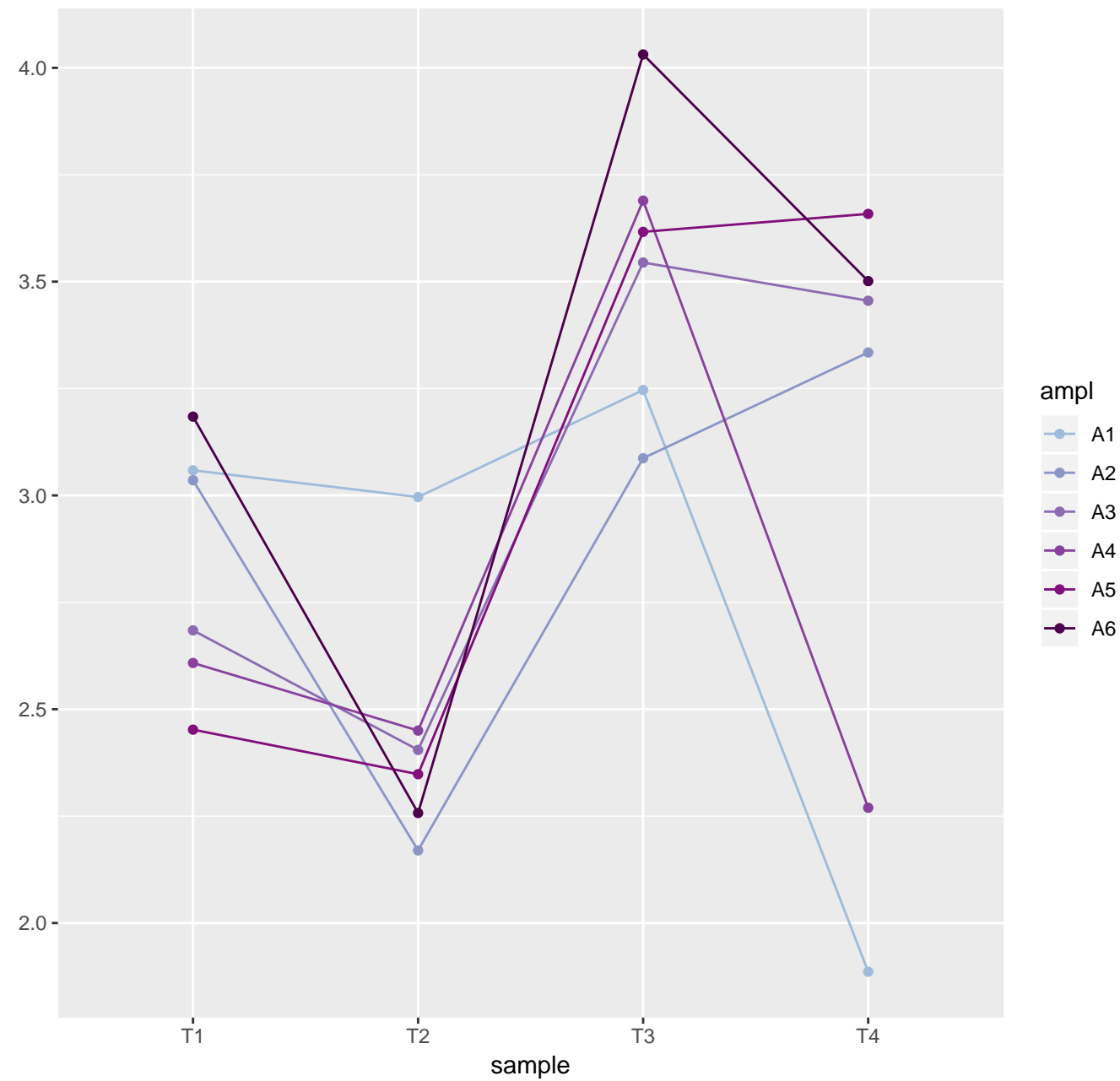




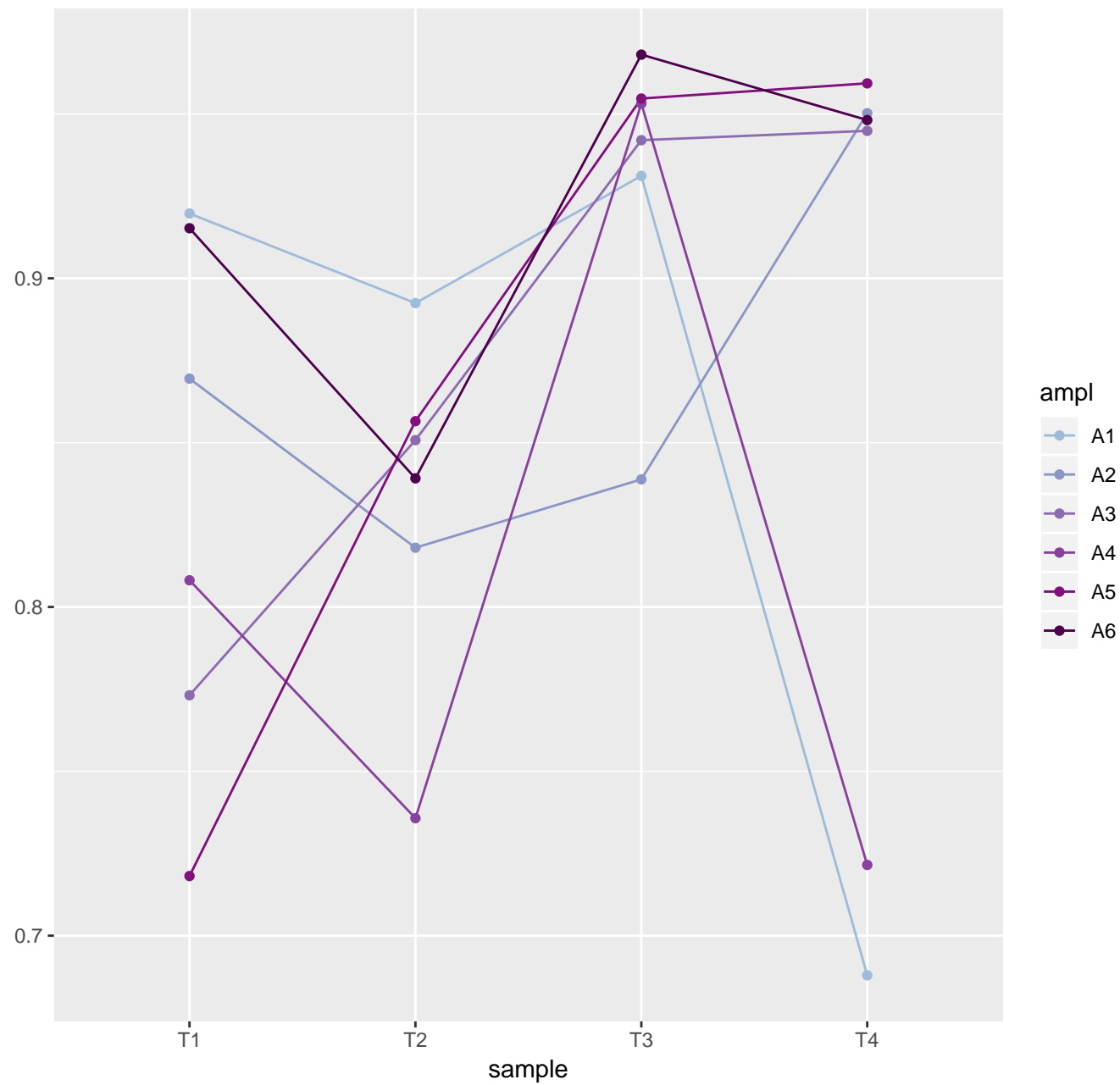
Havrda-Charvat estimator_0



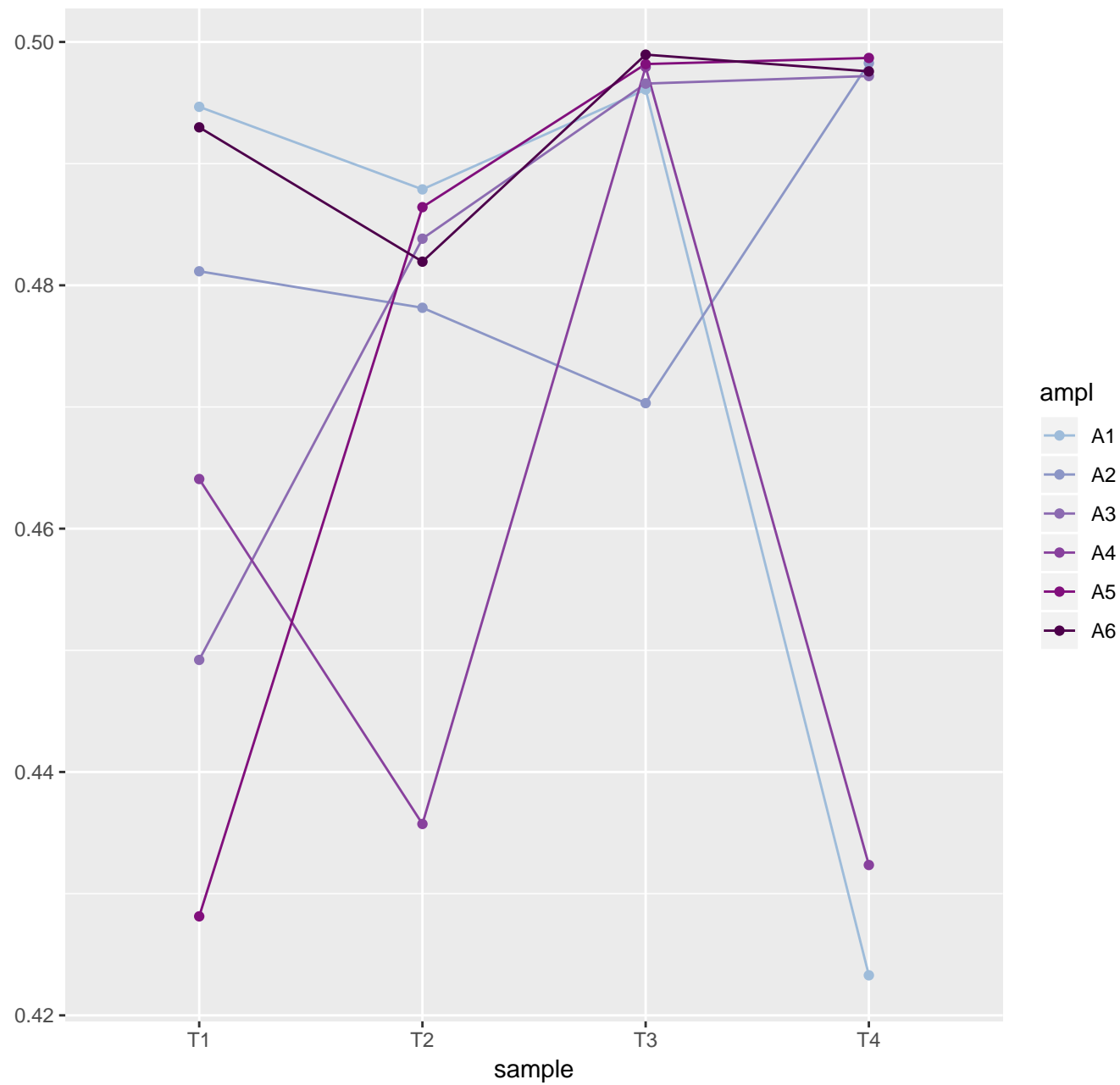
Havrda-Charvat estimator_1



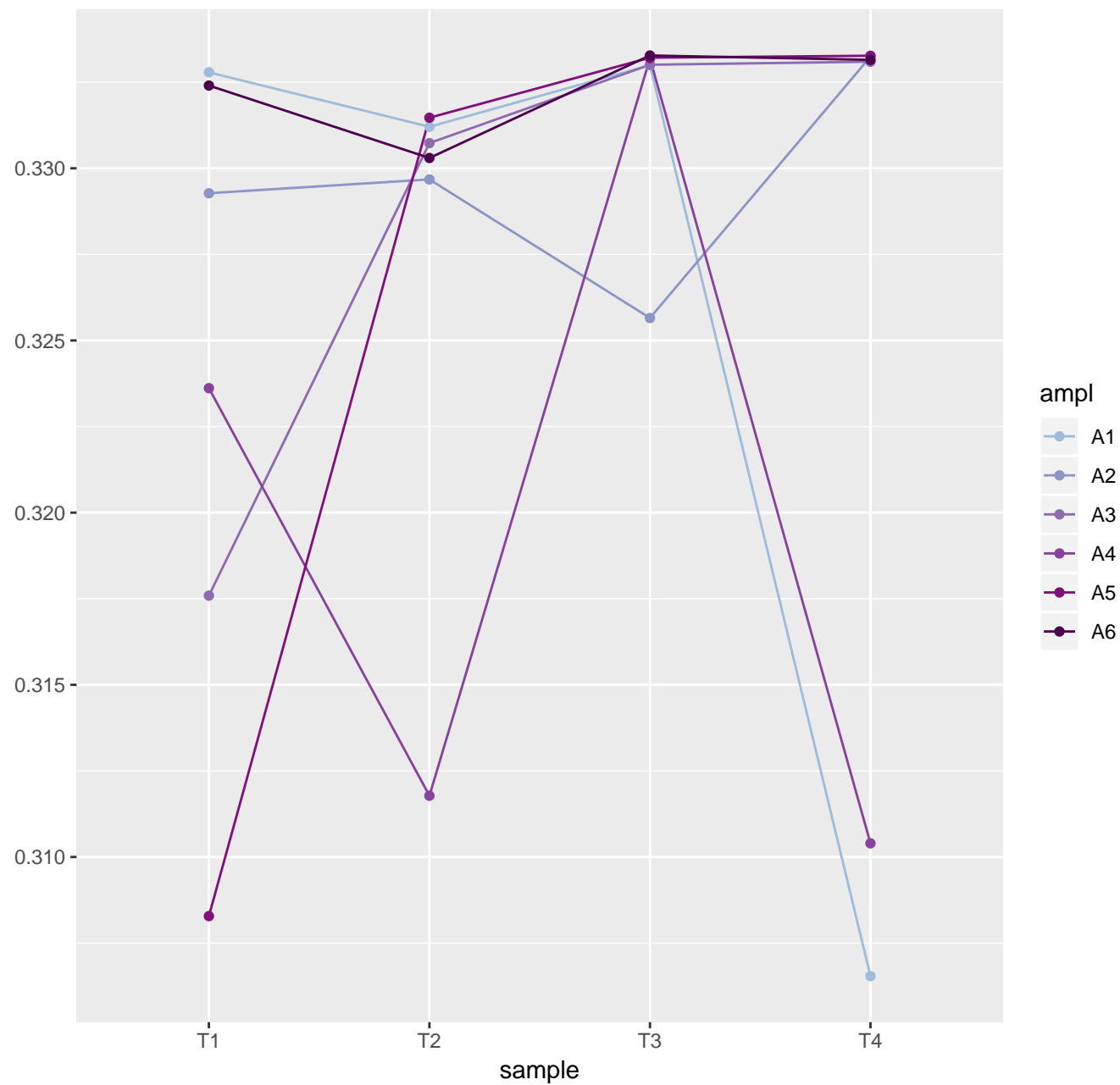
Havrda-Charvat estimator_2

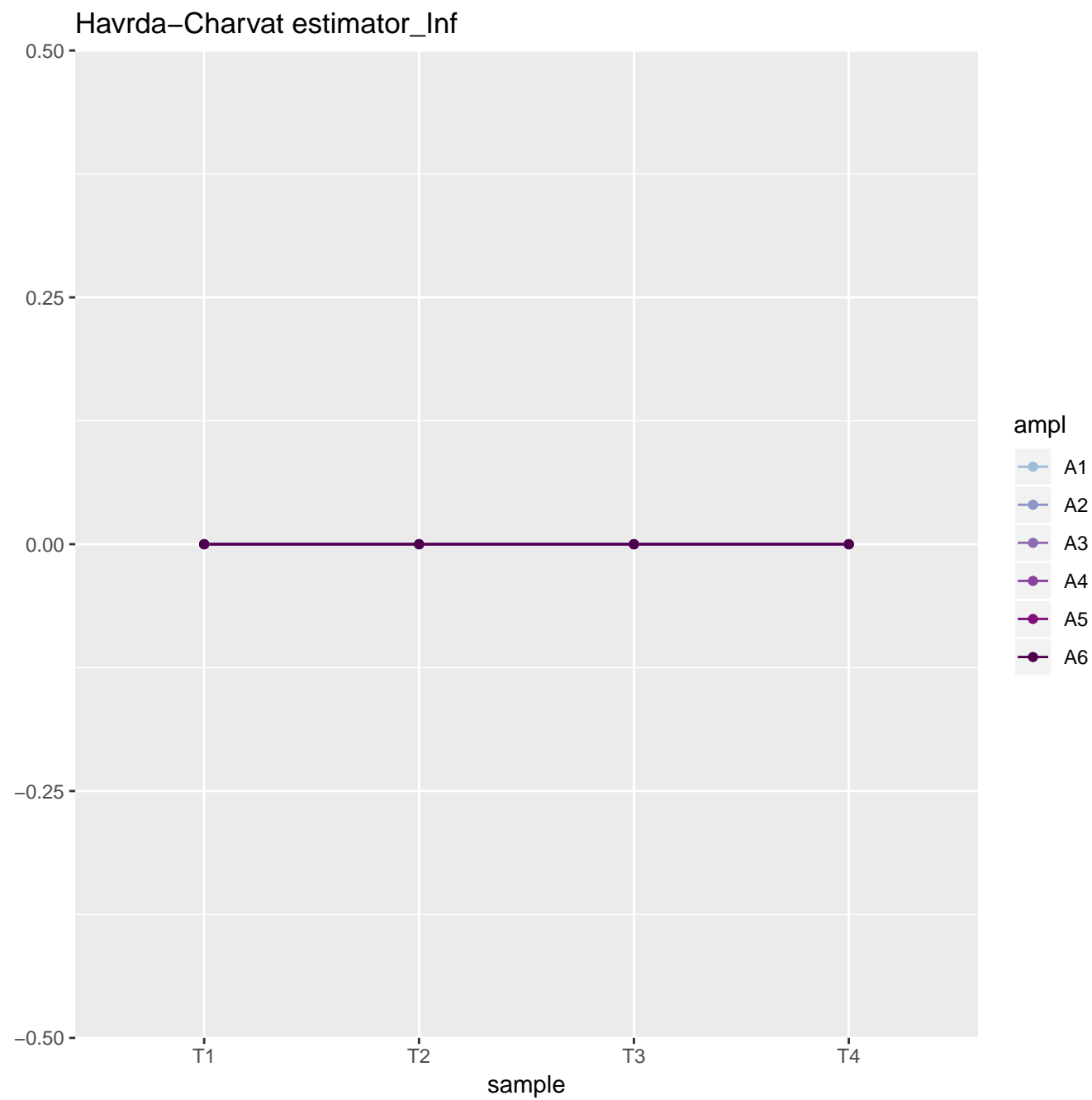


Havrda-Charvat estimator_3

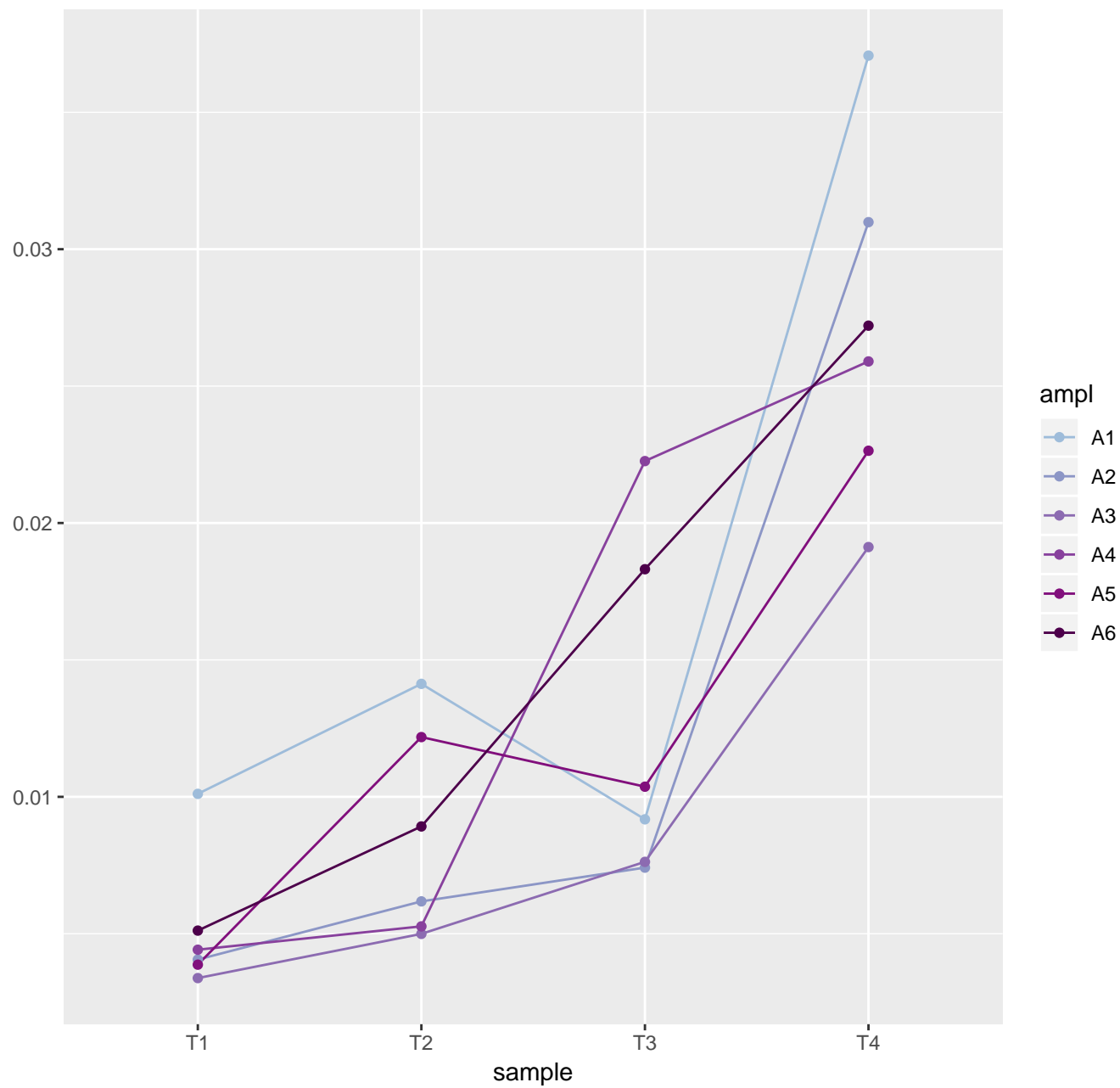


Havrda-Charvat estimator_4

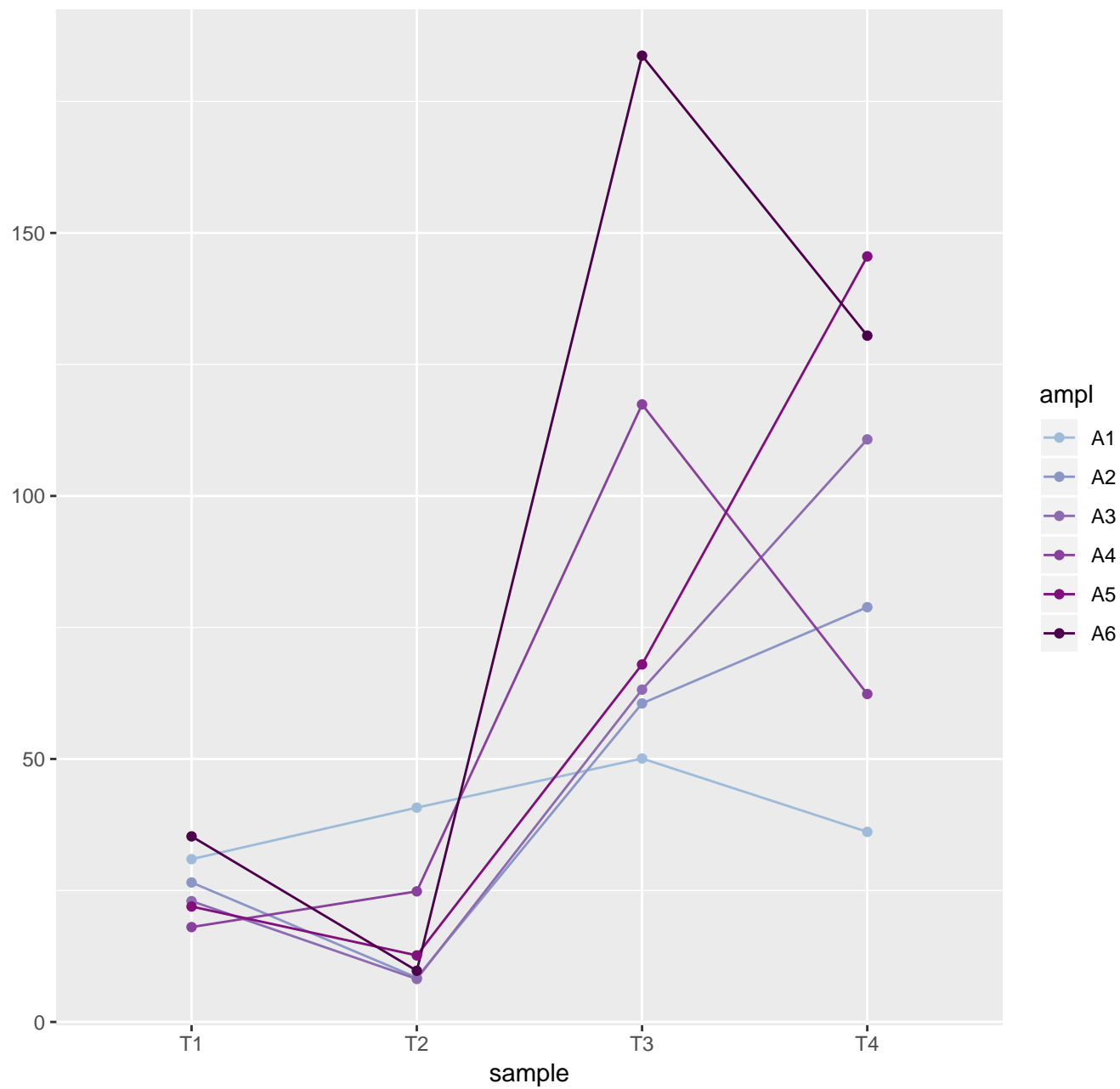




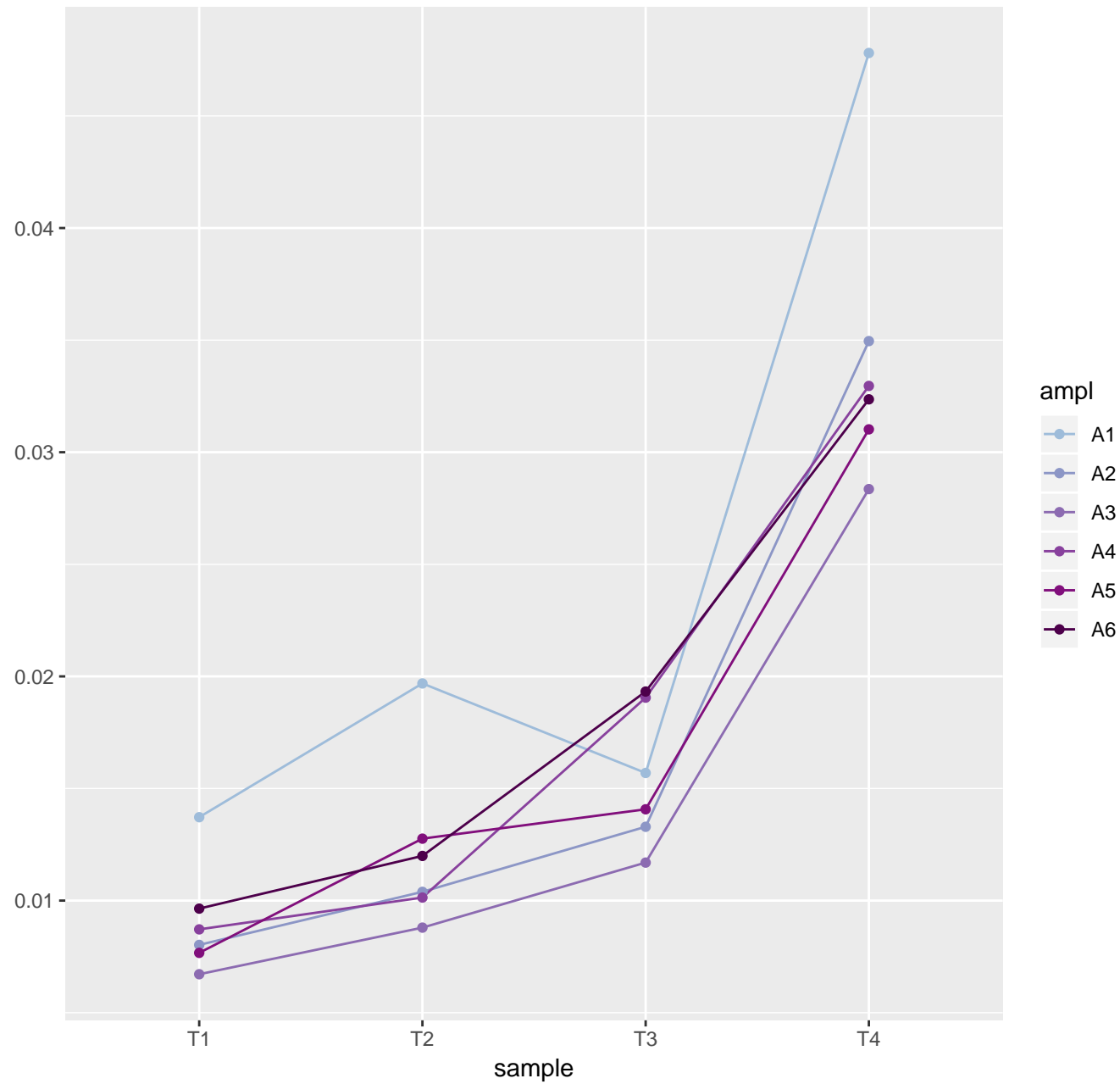
Average mutation frequency by entity



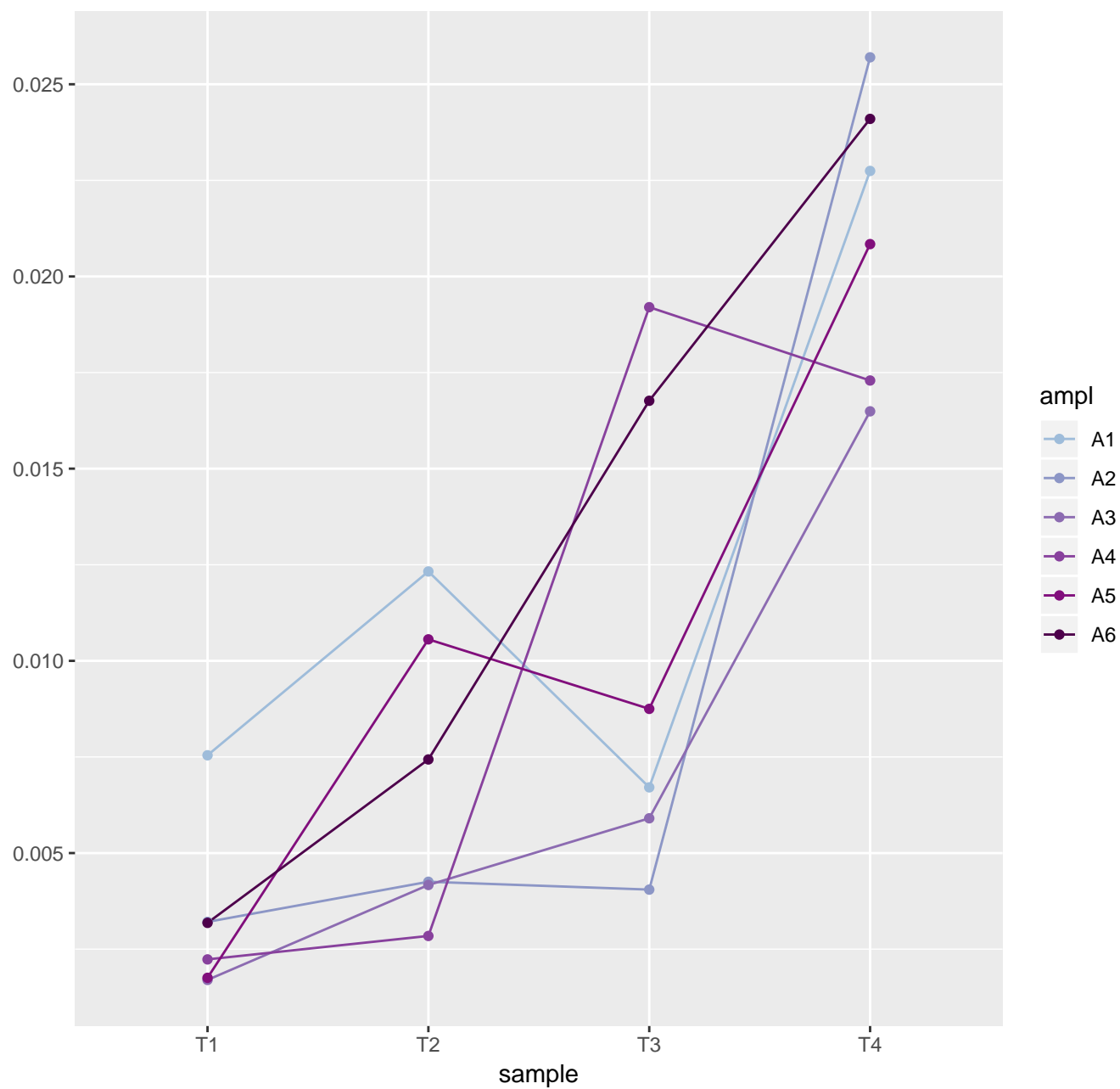
FAD



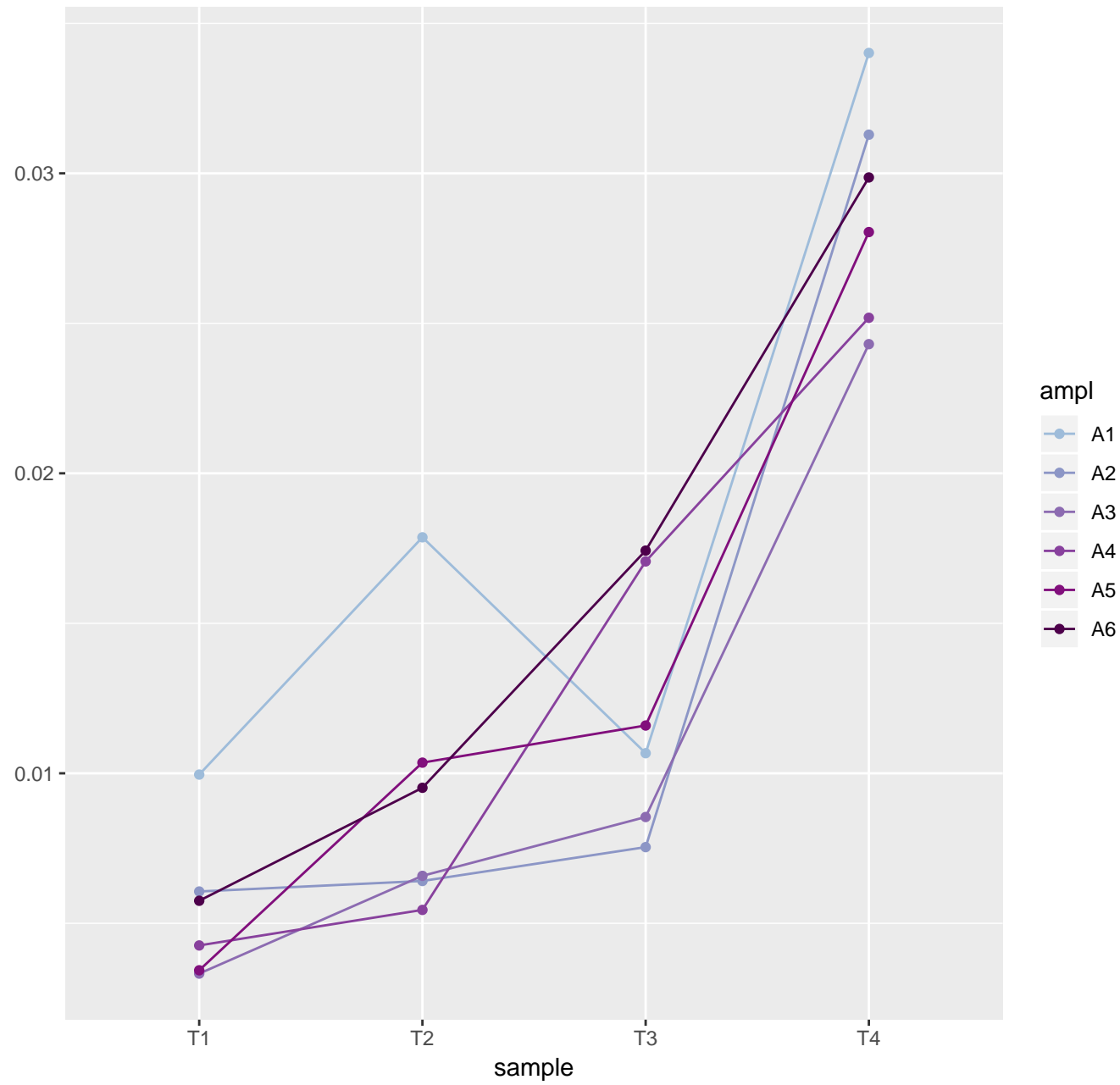
Nucleotide diversity by entity



Average mutation frequency by molecule



Nucleotide diversity



Rao entropy

